SEQUENCE LISTING

<110> della-Cioppa, Guy
    Erwin, Robert L.
    Fitzmaurice, Wayne P.
    Hanley, Kathleen M.
    Kumagai, Monto H.
    Lindbo, John A.
    McGee, David R.
    Padgett, Hal S.
    Pogue, Gregory P.

<120> METHOD OF DETERMINING THE FUNCTION OF
    NUCLEOTIDE SEQUENCES AND THE PROTEINS THEY ENCODE BY
    TRANSFECTING THE SAME INTO A HOST

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<140> 09/232,170
<141> 1999-01-15

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tttcctttac tcatacccaa acatgccgta cactcccttg cgggtggcct aaggactctt       360
gagttagagt atctcatgat gcaagttccg ttcggttctc tgacgtacga catcggtggt       420
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ctg ttt agc agg ctt ttt gcc aag aag gag atg cga att ctg atg gtt        96
Leu Phe Ser Arg Leu Phe Ala Lys Lys Glu Met Arg Ile Leu Met Val
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ggt ctt gat gct gct ggt aag acc aca atc ttg tac aag ctc aag ctc       144
Gly Leu Asp Ala Ala Gly Lys Thr Thr Ile Leu Tyr Lys Leu Lys Leu
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gga gag att gtc acc acc atc cct act att ggt ttc aat gtg gaa act       192
Gly Glu Ile Val Thr Thr Ile Pro Thr Ile Gly Phe Asn Val Glu Thr
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gtg gaa tac aag aac att agt ttc acc gtg tgg gat gtc ggg ggt cag       240
Val Glu Tyr Lys Asn Ile Ser Phe Thr Val Trp Asp Val Gly Gly Gln
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gac aag atc cgt ccc ttg tga gac act act tcc aga aca ctc aag gtc       288
Asp Lys Ile Arg Pro Leu  *  Asp Thr Thr Ser Arg Thr Leu Lys Val
        80                  85                  90

taa tct ttg ttg ttg ata gca atg aca gag aca gag ttg ttg agg ctc       336
 *  Ser Leu Leu Leu Ile Ala Met Thr Glu Thr Glu Leu Leu Arg Leu
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gag atg aac tcc aca gga tgc tga atg agg acg agc tgc gtg atg ctg       384
Glu Met Asn Ser Thr Gly Cys  *  Met Arg Thr Ser Cys Val Met Leu
        110                 115                 120

tgt tgc ttg tgt ttg cca aca agc aag atc ttc caa atg cta tga acg       432
Cys Cys Leu Cys Leu Pro Thr Ser Lys Ile Phe Gln Met Leu  *  Thr
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aag gtc tgg act ggc tct cca aca aca tcg ctg gca agg cat gat gag          576
Lys Val Trp Thr Gly Ser Pro Thr Thr Ser Leu Ala Arg His Asp Glu
                175             180             185

gga gaa att gcg ttg cat cga gat gat tct gtc tgc tgt gtt ggg atc          624
Gly Glu Ile Ala Leu His Arg Asp Asp Ser Val Cys Cys Val Gly Ile
            190             195             200

tct ctc tgt ctt gat gca aga gag att ata aat att atc tga acc ttt         672
Ser Leu Cys Leu Asp Ala Arg Glu Ile Ile Asn Ile Ile  *  Thr Phe
        205             210             215

ttg ctt ttt tgg gta tgt gaa tgt ttc tta ttg tgc aag tag atg gtc         720
Leu Leu Phe Trp Val Cys Glu Cys Phe Leu Leu Cys Lys  *  Met Val
        220             225             230

ttg tac cta aaa att tac tag aag aac cct ttt aaa tag ctt tcg tgt         768
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